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REMARKS

The amendment to the specification is made solely to insert the Sequence Listing and SEQ ID NOS. into the specification. No new matter is introduced. The accompanying declaration by the inventor confirms that the sequences in the Sequence Listing are the same as those incorporated by reference into the original specification by providing the GenBank® Accession Number for each sequence. A signed declaration will be filed under separate cover.

Applicant requests that claims 17-49 be examined together without restriction. Each of SEQ ID NOs:6, 7, and 8 relate to the human thymidylate synthetase gene. SEQ ID NO:6 corresponds to the coding region. SEQ ID NO:7 corresponds to the promoter region. SEQ ID NO:9 corresponds to the exons. Three SEQ ID NOs are required even though only one gene is referenced because the original specification identified the sequences by providing three GenBank® Accession Numbers.

Each of the nucleotides specified in the claims represents a single nulceotide polymorphism in the human thymidylate synthetase gene.

Attached is a marked-up version of the changes being made by the current amendment. Please apply any other charges or credits to Deposit Account No. 06-1050.

Respectfully submitted,

Date:

24 Sept 2061

Anita L. Meiklejohn, Ph.D.

Reg. No. 35,283

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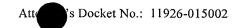
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Version with markings to show changes made

In the specification:

The paragraph beginning at page 1, line 8, has been amended as follows:

This application is a divisional of U.S. Application Serial No. 09/658,659, filed September 8, 2000, which is a CIP of Stanton, U.S. Application serial No. 09/596,033, filed June 15, 2000 entitled GENE SEQUENCE VARIACNES IN GENES RELATED TO FOLATE METABOLISM HAVING UTILITY IN DETERMINING THE TREATMENT OF DISEASE which is a CIP of Stanton, U.S. Application 09/357,743, filed July 20, 1999, entitled GENE SEQUENCE VARIACNES IN GENES RELATED TO FOLATE METABOLISM HAVING UTILITY IN DETERMINING THE TREATMENT OF DISEASE which is a CIP of Stanton, U.S. Application Serial No. 09/357,024, filed July 19, 1999, entitled GENE SEQUENCE VARIACNES IN GENES RELATED TO FOLATE METABOLISM HAVING UTILITY IN DETERMINING THE TREATMENT OF DISEASE, which claims the benefit of Stanton, U.S. Provisional Application 60/093,484, filed July 20, 1998, entitled GENE SEQUENCE VARIACNES IN GENES RELATED TO FOLATE METABOLISM HAVING UTILITY IN DETERMINING THE TREATMENT OF DISEASE, which are all hereby incorporated by reference in their entireties including drawings and tables.

Table 10 beginning at page 171 has been amended as follows:

Table 10

Va	ri	an	ce	Ta	d	le

Hugo	G	ID	OMIM	ID	VGX	Symbol	Description
7	/ariance	Start	Var	riance			
บ7333	38 U	173338	15657	70	GEN-	-69	Methionine
Synthase	(SEQ II	NO:1)					
_	194		(-2)	201)C>G	7	5	Ť
	284		(-1	11)C>T	7	5	ĭ
	1136)		742G>F	Ā	V2481	M
1252			858C>T		Silent	t	

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Filed : HEREWITH Page : 17 1334 940G>A D314N 1699 1305T>C Silent 3150 2756A>G D919G 3207 2813G>T S938I 3209 G939R 2815G>C 5444 5050C>A 3 ' 5551 5157G>A 3 ' 5573 5179C>T 3 ' 3 ' 5659 5265T>C 5678 3 ' 5284T>C 3 * 5874 5480C>T 5934 5540A>G 3 ' D78586 D78586 GEN-BR 114010 CAD PROTEIN (SEQ ID NO:2) 3434 3408C>T Silent 4313 4287T>C Silent 4799 4773A>G Silent 5255 5229C>T Silent 5455 5429G>A R18100 5507 5481T>C Silent 5810 5784C>T Silent 6128 6102C>T Silent 6626 6600C>T Silent 6686 6660C>T Silent U09178 U09178 274270 GEN-HA Dihydropyrimidine Dehydrogenase (SEQ ID NO:3) 166 85T>C C29R 577 496A>G M166V 638 557A>G Y186C 1708 1627A>G I543V 3432 3351T>C 3 ' 3 ' 3682 3601C>T 3730 3649G>A 3 ' 3 ' 3925 3844A>G 3937 3856T>C 3 **'** U19720 U19720 600424 GEN-I1 Folate Transporter (SLC19A1) (SEQ ID NO:4) 175 80G>A R27H 341 246C>G Silent 791 696C>T Silent 1067 972G>A Silent 1337 1242C>A Silent 1997 3 ' 1902T>C 3 **'** 2100 2005^2006insG 2582 3 ' 2487T>G

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2617	2522C>T	3'
2652	2557T>C	3'
		omo sapiens reduced
folate carrier (RFC1) g		
431	431A>G	Intron
441	431A>G	
498		Intron
579	498C>T	Intron
	579G>C	Intron
599	599G>C	Intron
X02308 X02308	188350 GEN-	KL Thymidylate
synthetase (SEQ ID NO:6		
1066	961T>C	3'
1136	1031A>G	3'
1497	1392T>A	3'
	88350 GEN-LUC	Thymidylate
synthase, promoter (SEQ	ID NO:7)	
276	276C>T	Intron
321	321T>C	Intron
452	452G>A	Intron
457	457^insC	Intron
491	491C>A	Intron
533	533T>C	Intron
624	624A>C	Intron
639	639A>G	Intron
655	655T>C	Intron
D00596 D00596	188350 GEN-L	
gene for thymidylate sy		
complete cds (SEQ ID NO		, 3, 1, 3, 3, 1,
701	701A>C	Intron
716	716A>G	Intron
732	732T>C	Intron
1293	1293A>G	Intron
1322	1322C>G	Intron
1379	1379T>C	Intron
1579		
1688	1590C>T	Intron
	1688C>G	Intron
2401	2401A>G	Intron
2429	2429G>A	Intron
2488	2488C>T	Intron
2594	2594G>T	Intron
2618	2618G>A	Intron
3083	3083G>A	Intron
3125	3125G>A	Intron
3212	3212C>T	Intron
3619	3619T>A	Intron

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0.00 **	0.00 = 0. 7	
3635	3635G>A	Intron
4256	4256G>A	Intron
4898	4898A>G	Intron
5006	5006C>T	Intron
5062	5062G>A	Intron
5167	5167G>A	Intron
11069	11069A>G	Intron
11238	11238C>T	Intron
11293	11293T>G	Intron
11422	11422T>C	Intron
11686	11686C>T	Intron
12598	12598T>C	Intron
13171	13171T>C	Intron
13298	13298G>A	Intron
13645	13645T>C	Intron
13751	13751C>A	Intron
13782	13782T>C	Intron
13806	13806T>C	Intron
13813	13813T>C	Intron
14479	14479A>G	Intron
14546	14546^insT	Intron
14585	14585C>T	Intron
14729	14729G>A	Intron
14787	14787C>T	Intron
14795	14795G>A	Intron
15041	15041T>C	Intron
15343	15343G>A	Intron
15449	15449G>A	Intron
15502	15502G>A	Intron
15545	15545C>T	Intron
15589	15589A>G	Intron
15769	15769C>T	3'
15839	15839A>G	3'
16148	16148G>A	3'
16198	16198T>G	3'
16202	16202G>T	Intron
X59618 X59618		I-M3 Ribonucleotide
reductase M2 polypeptide		
128	(-67)G>A	5 '
189	(-6)T>G	5'
524	330C>G	Silent
1399	1205T>A	3'
1464	1270G>A	3'
1636	1442C>T	3'
1738	1544C>T	3'
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2259 2065T>C S72487 GEN-3LD Thymidine S72487 131222 phosphorylase, partial (SEQ ID NO:10) 183 19G>A D7N 3 ' 319C>T 483 3 ' 437G>C 601 3 1 1135G>A 1299 GEN-LUB Thymidine M58602 M58602 131222 phosphorylase, promoter and genomic (SEQ ID NO:11) 3 ' 124C>T 124 3 ' 439G>A 439 3 ' 1044^insCT 1044 3 1 1331G>A 1331 1977 1977G>A Intron 2149G>A 2149 Intron 2467 2467A>G Intron 2634C>G Intron 2634 2975G>A 2975 Intron 3116G>T Intron 3116 3255 3255A>C Intron 3344 3344T>C Intron 4051C>A Intron 4051 4782G>A Intron 4782 5022T>C Intron 5022 5266 5266G>A Intron 5285 5285C>G Intron 5438 5438T>A Intron 5482C>T 5482 Intron 5629 5629G>A Intron 5648 5648C>T Intron 5731 5731G>A Intron M98045 M98045 136510 GEN-4C3 Homo sapiens folylpolyglutamate synthetase mRNA, complete cds (SEQ ID NO:12) Silent 732C>T 802 1747 1677G>T 3 ' 31 1900 1830T>C Human U24253 U24253 136510 GEN-LUE folylpolyglutamate synthetase (FPGS) gene, exons 5-11, and partial cds (SEQ ID NO:13) 1424 1424C>A Intron 1649 1649G>A Intron 2554 2554A>G Intron U24252 136510 GEN-LUF U24252 Folylpolyglutamate synthetase, promoter and exons 1-4 (SEQ ID NO:14)

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263	263A>G	Intron
266	266G>T	Intron
527	527C>G	Intron
1037	1037A>G	5 '
1139	1139G>A	Intron
1217	1217C>T	Intron
1647	1647C>T	Intron
1955	1955G>A	Intron
2017	2017G>A	Intron
2037	2037G>A	Intron
2189	2189A>G	Intron
2282	2282C>T	Intron
2309	2309A>G	Intron
U09806 U09806	236250 GEN-4E	FZ Human
methylenetetrahydrofol	ate reductase mRNA,	partial cds (SEQ ID
NO:15)		
120	120T>C	Silent
464		M155R
519	519C>T	Silent
668	668C>T	A223V
1059	1059T>C	Silent
1289	1289C>A	3'
1308	1308T>C	3'
1784	1784G>A	3'
AF061655 AF061655	123920 GEN-LUJ	Cytidine
deaminase, promoter (S		
575	575T>C	Intron
648	648T>C	Intron
771	771G>C	Intron
883	883G>A	Intron
941	941^insC	5 '
1051	1051A>C	K27Q